acid-polar amino acid-hydrophobic amino acid-D, wherein said peptide does not have toxin agonist activity, and wherein Xaa is an amino acid.

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(New) An isolated and purified peptide having an amino acid sequence  $Xaa_{(2)}KXaa_{(3)}TXaaQEXaaD$  wherein said peptide does not have toxin agonist activity, wherein Xaa is an amino acid.

(New) An isolated and purified peptide having an amino acid sequence Xaa<sub>(2)</sub>KKXaa<sub>(6)</sub>LD wherein said peptide does not have toxin agonist activity, wherein Xaa is an amino acid.

(New) An isolated an purified peptide having an amino acid sequence Xaa<sub>(2)</sub>-charged amino acid-Xaa<sub>(2)</sub>-hydrophobic amino acid-X-hydrophobic amino acid-polar amino acid-polar amino acid-hydrophobic amino acid-D, wherein said peptide does not have toxin agonist activity, and wherein Xaa is an amino acid.

## IN THE SPECIFICATION:

Please rewrite the following paragraphs as follows:

## On page 1, replace lines 11-14 as follows:

The U.S. Government has a nonexclusive, nontransferable, irrevocable paid-up license to practice or have practiced this invention for or on its behalf as provided for by the terms of Contract No. DAMD17-93-C-3108 awarded by the U.S. Department of the Army.

## On page 26, replace lines 1-11 as follows:

domain are conserved among all staphylococcal enterotoxins [Swaminathan et al. (1992) ibid.]. Indeed, the 150-161 domain of SEB is highly conserved among pyrogenic toxins in general, with 10/12 identities for SEA, SEC1, SEC2, and S. pyogenes exotoxin A (SPEA) and 9/12 for SEE [Bohach and Schlievert, Mol Gen Genet 209:5 (1987); Couch et al., J Bacteriol 170:2954 (1988);